<110> Burnham, Martin K. R.

<120> dexB

<130> GM10087

<150> 60/057,876
<151> 1997-09-02

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aag agt ttt atg gat agt aat gga gat gga gtt ggt gat ttg cca ggt   96
Lys Ser Phe Met Asp Ser Asn Gly Asp Gly Val Gly Asp Leu Pro Gly
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att acc agt aag ttg gac tat cta gct aag tta gga atc aca tcg att   144
Ile Thr Ser Lys Leu Asp Tyr Leu Ala Lys Leu Gly Ile Thr Ser Ile

tgg ctt tct ccc gtt tat gac agc cct atg gat gat aat ggc tac gat
192
Trp Leu Ser Pro Val Tyr Asp Ser Pro Met Asp Asp Asn Gly Tyr Asp
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att gct gat tat caa gcg att gcg gct att ttt gga acc atg gag gac
240
Ile Ala Asp Tyr Gln Ala Ile Ala Ala Ile Phe Gly Thr Met Glu Asp
     65   70   75   80

atg gat gaa ctg att gca gaa gct aag aag cgt gat atc cgt atc atc
288
Met Asp Glu Leu Ile Ala Glu Ala Lys Lys Arg Asp Ile Arg Ile Ile
                     85   90   95

atg gac ttg gtg gtc aat cat acc tcg gat gag cat gcc tgg ttt gta
336
Met Asp Leu Val Val Asn His Thr Ser Asp Glu His Ala Trp Phe Val
                 100   105   110

gag gcc tgt gaa aat cct aat agc cct gag cga gac tac tat atc tgg
384
Glu Ala Cys Glu Asn Pro Asn Ser Pro Glu Arg Asp Tyr Tyr Ile Trp
             115   120   125

cgc gat gaa ccc aat gac cta gat tct atc ttt agt ggg tct gct tgg
432
Arg Asp Glu Pro Asn Asp Leu Asp Ser Ile Phe Ser Gly Ser Ala Trp
         130   135   140

gaa tac gat gaa aag tca ggt caa tac tat ctc cac ttt ttc agc aag
480
Glu Tyr Asp Glu Lys Ser Gly Gln Tyr Tyr Leu His Phe Phe Ser Lys
145   150   155   160

aaa cag ccg gat ctc aac tgg gaa aat gaa aaa ctt cgc cag aaa att
528
Lys Gln Pro Asp Leu Asn Trp Glu Asn Glu Lys Leu Arg Gln Lys Ile

- 2 -

tat gag atg atg aac ttc tgg att gat aag ggt att ggt ggt ttc cgt
576
Tyr Glu Met Met Asn Phe Trp Ile Asp Lys Gly Ile Gly Gly Phe Arg
                180   185   190

atg gat gtt att gac atg att ggc aaa att cct gac gag aag gta gtc
624
Met Asp Val Ile Asp Met Ile Gly Lys Ile Pro Asp Glu Lys Val Val
            195   200   205

aat aat ggt cct atg ctc cat ccc tat ctc aag gaa atg aat cag gcg
672
Asn Asn Gly Pro Met Leu His Pro Tyr Leu Lys Glu Met Asn Gln Ala
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acc ttt gga gat aag gat ctc ttg aca gta ggg gag act tgg gga gca
720
Thr Phe Gly Asp Lys Asp Leu Leu Thr Val Gly Glu Thr Trp Gly Ala
225   230   235   240

acg cca gag att gcc aaa ctc tac tct gat cca aag ggg caa gaa ttg
768
Thr Pro Glu Ile Ala Lys Leu Tyr Ser Asp Pro Lys Gly Gln Glu Leu
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tct atg gtc ttc cag ttt gaa cat atc ggt ctt cag tat cag gaa ggt
816
Ser Met Val Phe Gln Phe Glu His Ile Gly Leu Gln Tyr Gln Glu Gly
                260   265   270

cag cct aaa tgg cac tat caa aaa gag ctg aat atc gct aag tta aaa
864
Gln Pro Lys Trp His Tyr Gln Lys Glu Leu Asn Ile Ala Lys Leu Lys
                275   280   285

gaa atc ttc aac aaa tgg cag aca gag tta gga gtt gag gac ggc tgg
912
Glu Ile Phe Asn Lys Trp Gln Thr Glu Leu Gly Val Glu Asp Gly Trp

-3-

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960
Asn Ser Leu Phe Trp Asn Asn His Asp Leu Pro Arg Ile Val Ser Ile
305   310   315   320

tgg gga aat gac caa gaa tac cgc gaa aaa tct gcc aaa gcc ttt gca
1008
Trp Gly Asn Asp Gln Glu Tyr Arg Glu Lys Ser Ala Lys Ala Phe Ala
                 325   330   335

atc ttg ctt cat ctt atg aga gga act cct tat atc tac caa ggt gag
1056
Ile Leu Leu His Leu Met Arg Gly Thr Pro Tyr Ile Tyr Gln Gly Glu
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gag att ggg atg acc aac tat ccg ttt gaa aca ctg gat caa gta gaa
1104
Glu Ile Gly Met Thr Asn Tyr Pro Phe Glu Thr Leu Asp Gln Val Glu
         355   360   365

gat att gaa tct ctc aac tat gcg cgt gag gct ctt gaa aaa ggt gtt
1152
Asp Ile Glu Ser Leu Asn Tyr Ala Arg Glu Ala Leu Glu Lys Gly Val
     370   375   380

ccg atg caa gaa atc atg gac agt atc cgt gtt att gga cgt gac aat
1200
Pro Met Gln Glu Ile Met Asp Ser Ile Arg Val Ile Gly Arg Asp Asn
385   390   395   400

gcc cgt acc cct atg caa tgg gac gag agc aaa aac gct ggt ttc tca
1248
Ala Arg Thr Pro Met Gln Trp Asp Glu Ser Lys Asn Ala Gly Phe Ser
                 405   410   415

aca ggt caa cct tgg ttg gca gtt aat cca aat tac gag atg atc aac
1296
Thr Gly Gln Pro Trp Leu Ala Val Asn Pro Asn Tyr Glu Met Ile Asn

gtc caa gaa gcg ctg gca aat cca gat tct att ttc tat acc tat cag
1344
Val Gln Glu Ala Leu Ala Asn Pro Asp Ser Ile Phe Tyr Thr Tyr Gln
      435   440   445

aaa ctg gtc caa att cgc aag gag aat agt tgg cta att cga gct gac
1392
Lys Leu Val Gln Ile Arg Lys Glu Asn Ser Trp Leu Ile Arg Ala Asp
      450   455   460

ttt gaa ttg ctt gat acg gct gat aag gtc ttt gct tat ata cgt aag
1440
Phe Glu Leu Leu Asp Thr Ala Asp Lys Val Phe Ala Tyr Ile Arg Lys
465   470   475   480

gat ggc gac cgt cgc ttc cta gtt gtg gct aac ttg tcc aat gaa gag
1488
Asp Gly Asp Arg Arg Phe Leu Val Val Ala Asn Leu Ser Asn Glu Glu
              485   490   495

caa gac ttg aca gta gaa gga aaa gtc aaa tct gtc ttg att gaa aac
1536
Gln Asp Leu Thr Val Glu Gly Lys Val Lys Ser Val Leu Ile Glu Asn
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acc cta gct caa gaa gtc ttt gaa aaa caa atc tta gtt cca tgg gat
1584
Thr Leu Ala Gln Glu Val Phe Glu Lys Gln Ile Leu Val Pro Trp Asp
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            35                  40                  45
Trp Leu Ser Pro Val Tyr Asp Ser Pro Met Asp Asp Asn Gly Tyr Asp
        50                  55                  60
Ile Ala Asp Tyr Gln Ala Ile Ala Ala Ile Phe Gly Thr Met Glu Asp
65                  70                  75                  80
Met Asp Glu Leu Ile Ala Glu Ala Lys Lys Arg Asp Ile Arg Ile Ile
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Met Asp Leu Val Val Asn His Thr Ser Asp Glu His Ala Trp Phe Val
                100                 105                 110
Glu Ala Cys Glu Asn Pro Asn Ser Pro Glu Arg Asp Tyr Tyr Ile Trp
            115                 120                 125
Arg Asp Glu Pro Asn Asp Leu Asp Ser Ile Phe Ser Gly Ser Ala Trp
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Glu Tyr Asp Glu Lys Ser Gly Gln Tyr Tyr Leu His Phe Phe Ser Lys
145                 150                 155                 160
Lys Gln Pro Asp Leu Asn Trp Glu Asn Glu Lys Leu Arg Gln Lys Ile
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Tyr Glu Met Met Asn Phe Trp Ile Asp Lys Gly Ile Gly Gly Phe Arg
            180                 185                 190
Met Asp Val Ile Asp Met Ile Gly Lys Ile Pro Asp Glu Lys Val Val
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Asn Asn Gly Pro Met Leu His Pro Tyr Leu Lys Glu Met Asn Gln Ala
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Thr Phe Gly Asp Lys Asp Leu Leu Thr Val Gly Glu Thr Trp Gly Ala
225                 230                 235                 240
Thr Pro Glu Ile Ala Lys Leu Tyr Ser Asp Pro Lys Gly Gln Glu Leu
            245                 250                 255
Ser Met Val Phe Gln Phe Glu His Ile Gly Leu Gln Tyr Gln Glu Gly
            260                 265                 270
Gln Pro Lys Trp His Tyr Gln Lys Glu Leu Asn Ile Ala Lys Leu Lys
        275                 280                 285
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Glu Ile Phe Asn Lys Trp Gln Thr Glu Leu Gly Val Glu Asp Gly Trp
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Asn Ser Leu Phe Trp Asn Asn His Asp Leu Pro Arg Ile Val Ser Ile
305   310   315   320
Trp Gly Asn Asp Gln Glu Tyr Arg Glu Lys Ser Ala Lys Ala Phe Ala
            325   330   335
Ile Leu Leu His Leu Met Arg Gly Thr Pro Tyr Ile Tyr Gln Gly Glu
            340   345   350
Glu Ile Gly Met Thr Asn Tyr Pro Phe Glu Thr Leu Asp Gln Val Glu
    355   360   365
Asp Ile Glu Ser Leu Asn Tyr Ala Arg Glu Ala Leu Glu Lys Gly Val
    370   375   380
Pro Met Gln Glu Ile Met Asp Ser Ile Arg Val Ile Gly Arg Asp Asn
385   390   395   400
Ala Arg Thr Pro Met Gln Trp Asp Glu Ser Lys Asn Ala Gly Phe Ser
            405   410   415
Thr Gly Gln Pro Trp Leu Ala Val Asn Pro Asn Tyr Glu Met Ile Asn
            420   425   430
Val Gln Glu Ala Leu Ala Asn Pro Asp Ser Ile Phe Tyr Thr Tyr Gln
    435   440   445
Lys Leu Val Gln Ile Arg Lys Glu Asn Ser Trp Leu Ile Arg Ala Asp
    450   455   460
Phe Glu Leu Leu Asp Thr Ala Asp Lys Val Phe Ala Tyr Ile Arg Lys
465   470   475   480
Asp Gly Asp Arg Arg Phe Leu Val Val Ala Asn Leu Ser Asn Glu Glu
            485   490   495
Gln Asp Leu Thr Val Glu Gly Lys Val Lys Ser Val Leu Ile Glu Asn
            500   505   510
Thr Leu Ala Gln Glu Val Phe Glu Lys Gln Ile Leu Val Pro Trp Asp
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